

# NEURON for multiscale simulations: reaction-diffusion and electrophysiology

RA McDougal<sup>1</sup>, A Bulanova<sup>1</sup>, M.N.I. Patoary<sup>2</sup>, C Tropper<sup>2</sup>, ML Hines<sup>1</sup>, WW Lytton<sup>3,4</sup> 1 Yale University, 2 McGill University, 3 SUNY Downstate, 4 Kings County Hospital



#### Introduction

- NEURON is widely used for simulating electrophysiology.
- NEURON 7.3 included a module for reaction-diffusion simulation, to facilitate the study of chemical-electrical interactions.
- Current work includes expanding reaction-diffusion capabilities and interfacing different techniques.
- Model specification is independent of discretization or solver.

#### **Plugins**

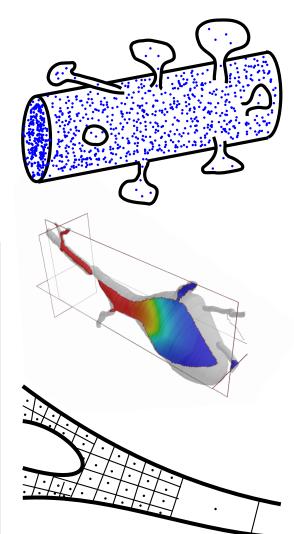
To support functionality not yet implemented in NEURON and domain-specific solvers, we are developing a plugin framework. Plugins must provide:

- sol.init(species, reactions)
- $\bullet$  sol.advance(dt)
- sol.set\_rate\_of\_change(nodelist, values)
- sol.transfer\_states\_to\_neuron(which=None)
- sol.transfer\_states\_from\_neuron()
- sol.supported\_sim()

Activate the solver via:

• rxd.set\_solver(sol)

## Interfaces



Small volumes are best modeled stochastically;

changes in geometry are best modeled with

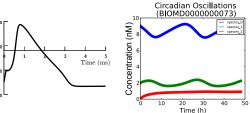
Difficulty: the right choice of 1D vs 3D or

deterministic vs stochastic may be different

a fine, high-dimensional mesh.

for different species.

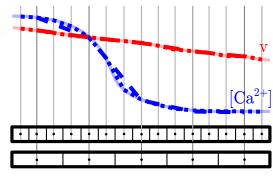
Time



Timescales may vary wildly within the same

In fixed step mode, NEURON currently offers the option of only updating the reaction-diffusion on some fraction of time steps.

#### Space



A fine discretization helps capture changes in chemical concentrations over short distances; voltage typically takes large distances to vary significantly.

Challenges: Develop a clear and concise specification. Develop rules for branch points.

### Acknowledgments

Supported by R01MH086638, 2T15LM007056.

## Model Specification

Where?



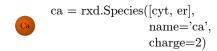
r = rxd.Region(apicals, geometry=???)



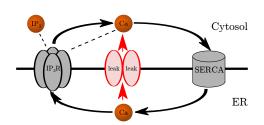




#### Who?



#### What?



leak = rxd.MultiCompartmentReaction(  $ca[er] \Leftrightarrow ca[cyt], kf, kb,$ membrane=ermem)

McDougal et al. Front Neuroinf. 2013.